	RF Errors Corrected by the STIC stems Branch
1 }	dumber: 09/751, 299 A Edited by:
	Changed a file from non-ASCII to ASCII
	Changed the margins in cases where the sequence text was 'wrapped' dewrond the margins in cases where the sequence text was 'wrapped'
	Edited a format error in the Current Application Data section, specifically:
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for *Current Application Data*.
	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
	Inserted colons after headings/subheadings. Headings edited included:
	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file page numbers throughout text; other invalid text, such as
	Inserted mandatory headings, specifically: <220>, sequences # 2, 4
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
	Corrected an error in the Number of Sequences field, specifically:
_	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
	Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error
	due to a Patentin bug). Sequences corrected:

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

#### OIPE

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/09/751,299A**DATE: 06/29/2001
TIME: 14:23:08

Input Set : A:\DIVER1440-2final.txt

Output Set: N:\CRF3\06292001\I751299A.raw

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3 <110> APPLICANT: Madden, Mark
         Weiner, David P.
         Chaplin, Jennifer A.
 7 <120> TITLE OF INVENTION: METHODS FOR PRODUCING ENANTIOMERICALLY PURE
         ALPHA-SUBSTITUTED CARBOXYLIC ACIDS
10 <130> FILE REFERENCE: DIVER1440-2
12 <140> CURRENT APPLICATION NUMBER: US 09/751,299A
13 <141> CURRENT FILING DATE: 2000-12-28
15 <150> PRIOR APPLICATION NUMBER: 60/254,414
16 <151> PRIOR FILING DATE: 2000-12-07
18 <150> PRIOR APPLICATION NUMBER: 60/173,609
19 <151> PRIOR FILING DATE: 1999-12-29
21 <160> NUMBER OF SEQ ID NOS: 4
23 <170> SOFTWARE: PatentIn Ver. 2.1
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34 <220> FEATURE:
35 <221> NAME/KEY: CDS
36 <222> LOCATION: (1)..(1041)
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41
                     5
                                        10
43 ccg gtg ttc ctc gat ctc gac cgc aca gtc gag aaa gcg atc ggc ctg
                                                                      96
44 Pro Val Phe Leu Asp Leu Asp Arg Thr Val Glu Lys Ala Ile Gly Leu
                20
                                    25
47 atc gag cag gcg gcc aag cag gac gtg cgc ctg atc gca ttc cca gag
                                                                      144
48 Ile Glu Gln Ala Ala Lys Gln Asp Val Arg Leu Ile Ala Phe Pro Glu
            35
                                40
51 act tgg att ccc ggc tat ccc ttt tgg ata tgg ctg ggc gcg ccg gct
52 Thr Trp Ile Pro Gly Tyr Pro Phe Trp Ile Trp Leu Gly Ala Pro Ala
        50
55 tgg ggc atg cgc ttc gtc cag cgc tat ttc gag aat tcg ctc gtg cgc
56 Trp Gly Met Arg Phe Val Gln Arg Tyr Phe Glu Asn Ser Leu Val Arg
                        70
                                             75
59 ggc agc aag cag tgg cag gcc ctg gcg gat gcg gcc cgc cgc cac ggc
                                                                      288
60 Gly Ser Lys Gln Trp Gln Ala Leu Ala Asp Ala Ala Arg Arg His Gly
61
                    85
                                         90
63 atg cat gtc gtg gcc ggc tat agc gag cgc gcg ggc ggc agc ctc tat
                                                                      336
64 Met His Val Val Ala Gly Tyr Ser Glu Arg Ala Gly Gly Ser Leu Tyr
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67 atg ggc cag gcg atc ttc ggc ccc gat ggc gat ctg atc gcc gcg cgc
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Input Set : A:\DIVER1440-2final.txt
Output Set: N:\CRF3\06292001\1751299A.raw

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71	cgc as Arg Ly	ag cto	aag		Thr		gcg			Thr		ttc				432
76 77	gac go Asp Gi 145	ly Ser	His	Leu	Ala 150	Val	His	Asp	Thr	Ala 155	Ile	Gly	Arg	Leu	Gly 160	480
80 81	gcg ct Ala Le	eu Cys	Cys	Trp 165	Glu	His	Ile	Gln	Pro 170	Leu	Ser	Lys	Tyr	Ala 175	Met	528
	tac go Tyr A															576
	ctc ta		Gly													624
	gca ac Ala Se 2:				Ala					Cys						672
96	tgc go Cys Al 225															720
	gac aa Asp 1				Leu					Gly					Phe	768
	Gly I			/ Arg					ı Pro					Glu		816
	gga o Gly 1		u Val					Let					Let			864
	gcg d Ala A		_	_			His					Asp	_	_		912
116	ctg of Leu l					Pro					Val					960
	gca 1 Ala 1				Asn					/ Asp					ı Arg	1008
124 125	gtg ( Val V ( <210)	Val Al	a Glu 340	ı Ser	-				a Gln			•				1041
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Input Set : A:\DIVER1440-2final.txt
Output Set: N:\CRF3\06292001\I751299A.raw

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Input Set : A:\DIVER1440-2final.txt
Output Set: N:\CRF3\06292001\I751299A.raw

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188 <220> FEATURE:														
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192 <220> FEATURE:														
193 <221> NAME/KEY: CDS														
194 <222> LOCATION: (1)(1014)														
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199 1 5 10	15													
201 atg gat ttg gag gcg acg gtg gac aaa acc att gag ttg at														
202 Met Asp Leu Glu Ala Thr Val Asp Lys Thr Ile Glu Leu Me														
`	30													
205 gca gca cgt aat aat gct cgt ctg atc gcc ttt ccg gaa ac														
206 Ala Ala Arg Asn Asn Ala Arg Leu Ile Ala Phe Pro Glu Th	nr Trp Ile													
207 35 40 45														
209 cca ggc tac cca tgg ttt ctt tgg ctt gac tca cca gca tg														
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221 acc ctg ggg atg agt gaa cgg gtc ggt ggc acc ctt tac at														
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227 115 120 125	ту пуз пец													
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230 Lys Pro Thr Phe Val Glu Arg Thr Leu Phe Gly Glu Gly As	2 3													
231 130 135 140														
233 tcg cta gcg gtt ttc gag acg tct gtt gga agg ctg ggt g	gc tta tgc 480													
234 Ser Leu Ala Val Phe Glu Thr Ser Val Gly Arg Leu Gly Gl														
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243 180 185 19														
245 aat gcg gcg aaa gcc ctg ggg cct gat gtc aat gta gcg gc	cc tct cga 624													
246 Asn Ala Ala Lys Ala Leu Gly Pro Asp Val Asn Val Ala Al	la Ser Arg													
247 195 200 . 205	•													
249 atc tat gcc gtt gaa ggg caa tgc ttc gta cta gcg tcg tg														
250 Ile Tyr Ala Val Glu Gly Gln Cys Phe Val Leu Ala Ser Cy	· Ala Iou													

Input Set : A:\DIVER1440-2final.txt
Output Set: N:\CRF3\06292001\I751299A.raw

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257	gcg '	ttq	ctt	ctq	gct	gat	gat	qqa	cac	tca	cqt	atc	ata	aga	cct	αat	768
	Ālā																
259					245	-	_	-		250.	,				255	[	
	. ggt	aat	σac	tta	atc	aca	cct	ctt	acc	gaa	aat	σaa	gag	aat		ctc	816
	Gly																•
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	Tyr																
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	) Asp		_					_					_	_			712
271		290		O <sub>T</sub>		- 1 -	295	••••		Пор	110	300	9	Lou	200	110	•
	gat		adc	cct	aaa	tta		ata	att	maa	att		aat	aat	ctt	cat	960
	Asp																500
	305	**** 9	501	110	Lyo	310	110	v a.i.	·uı	OIU	315	Olu	O <sub>1</sub> y	1100	пси	320	
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	Pro																1000
279		ı yı	nıα	пси	325	цуз	ΑΙα	561	Oiu	330	Gry	ALG	GIII	пси	335	Giu	
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289 289	<223	3> o:	rhèr	INF		CION:	Des	scrip	otion	n of	Unkr	nown	Orga	nism	n: Ok	otaine	d from an
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289 289 293 293 294 295 296	<pre> &lt;223 2 &lt;400 3 Met</pre>	3> Oʻ 0> SI Lys Asp	THÈR EQUEN Glu Leu	INFONCE: Ala Glu 20	4 Ile 5 Ala	Lys Thr	Val Val	Ala Asp	Cys Lys 25	Val 10 Thr	Gln Ile	Ala Glu	Ala Leu	Pro Met 30	Ile 15 Glu	Tyr Glu	d from an
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289 289 292 293 294 295 296 297 298	<pre>&lt;223 2 &lt;400 3 Met</pre>	3> On D> SI Lys Asp Ala Gly	THÈR EQUEN Glu Leu Arg 35	INFO NCE: Ala Glu 20 Asn	4 Ile 5 Ala Asn	Lys Thr Ala	Val Val Arg Leu	Ala Asp Leu 40	Cys Lys 25 Ile	Val 10 Thr	Gln Ile Phe	Ala Glu Pro Pro	Ala Leu Glu 45	Pro Met 30 Thr	Ile 15 Glu Trp	Tyr Glu Ile	d from an
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289 299 299 299 299 299 299 300 301 302 303 305 306	<pre></pre>	3> OT D> SI Lys Asp Ala Gly 50 Phe Ala Leu	THÈR THÈR Glu Leu Arg 35 Tyr Val Lys Gly Ile	INFO NCE: Ala Glu 20 Asn Pro Arg Arg Met 100	4 Ile 5 Ala Asn Trp Gln Ile 85 Ser	Lys Thr Ala Phe Tyr 70 Ser Glu	Val Val Arg Leu 55 His Asp Arg	Ala Asp Leu 40 Trp Glu Ala Val Asp	Cys Lys 25 Ile Leu Asn Ala Gly 105	Val 10 Thr Ala Asp Ser Lys 90 Gly	Gln Ile Phe Ser Leu 75 Arg	Ala Glu Pro Pro 60 Glu Leu Leu	Ala Leu Glu 45 Ala Leu Gly Tyr	Pro Met 30 Thr Trp Asp Ile Ile 110	Ile 15 Glu Trp Ala Gly Met 95 Ser	Tyr Glu Ile Met Pro 80 Val Gln	d from an
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289 299 299 299 299 299 299 300 301 303 304 305 306	<pre>  &lt;22:    &lt;400    Met         1    Met         Ala         Pro         Gln         65         Gln         Thr         Trp         Lys</pre>	3> OT D> SI Lys Asp Ala Gly 50 Phe Ala Leu	THÈR THÈR EQUEN Glu Leu Arg 35 Tyr Val Lys Gly Ile 115	INFO NCE: Ala Glu 20 Asn Pro Arg Arg Met 100 Gly	4 Ile 5 Ala Asn Trp Gln Ile 85 Ser Asp	Lys Thr Ala Phe Tyr 70 Ser Glu Asn	Val Val Arg Leu 55 His Asp Arg Gly	Ala Asp Leu 40 Trp Glu Ala Val Asp 120	Cys Lys 25 Ile Leu Asn Ala Gly 105 Thr	Val 10 Thr Ala Asp Ser Lys 90 Gly Ile	Gln Ile Phe Ser Leu 75 Arg Thr	Ala Glu Pro 60 Glu Leu Leu Ala	Ala Leu Glu 45 Ala Leu Gly Tyr Arg 125	Pro Met 30 Thr Trp Asp Ile Ile 110 Arg	Ile 15 Glu Trp Ala Gly Met 95 Ser Lys	Tyr Glu Ile Met Pro 80 Val Gln Leu	d from an

W-->

### VERIFICATION SUMMARY

PATENT APPLICATION: US/09/751,299A

DATE: 06/29/2001 TIME: 14:23:09

Input Set : A:\DIVER1440-2final.txt

Output Set: N:\CRF3\06292001\I751299A.raw

L:132 M:258 W: Mandatory Feature missing, <220> FEATURE: L:289 M:258 W: Mandatory Feature missing, <220> FEATURE:

STATISTICS SUMMARY

DATE: 06/29/2001 PATENT APPLICATION: US/09/751,299A TIME: 14:23:09

Input Set : A:\DIVER1440-2final.txt

Output Set: N:\CRF3\06292001\I751299A.raw

Application Serial Number: US/09/751,299A

Alpha or Numeric: Numeric

Application Class:

Application File Date: 12-28-2000

Art Unit: OIPE

Software Application: PatentIn Total Number of Sequences: 4

Total Nucleotides: 2055 Total Amino Acids: 683 Number of Errors: 0 Number of Warnings: 2 Number of Corrections: 0

#### MESSAGE SUMMARY

258 W: 2 (Mandatory Feature missing)

DATE: 05/31/2001

TIME: 17:55:10

### OIPE

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Does Not Comply
                                                                      Corrected Diskette Needed
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                     Output Set: C:\CRF3\05312001\I751299.raw
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              Weiner, David P.
              Chaplin, Jennifer A.
      7 <120> TITLE OF INVENTION: METHODS FOR PRODUCING ENANTIOMERICALLY PURE
              ALPHA-SUBSTITUTED CARBOXYLIC ACIDS
     10 <130> FILE REFERENCE: DIVER1440-2
     12 <140> CURRENT APPLICATION NUMBER: US 09/751,299
C--> 13 <141> CURRENT FILING DATE: 2001-05-01
     15 <150> PRIOR APPLICATION NUMBER: 60/254,414
     16 <151> PRIOR FILING DATE: 2000-12-07
     18 <150> PRIOR APPLICATION NUMBER: 60/173,609
     19 <151> PRIOR FILING DATE: 1999-12-29
     21 <160> NUMBER OF SEQ ID NOS: 4
     23 <170> SOFTWARE: PatentIn Ver. 2.1
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     27 <212> TYPE: DNA
     28 <213> ORGANISM: Unknown Organism
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     35 <221> NAME/KEY: CDS
     36 <222> LOCATION: (1)..(1041)
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                                                                           96
     43 ccg gtg ttc ctc gat ctc gac cgc aca gtc gag aaa gcg atc ggc ctg
     44 Pro Val Phe Leu Asp Leu Asp Arg Thr Val Glu Lys Ala Ile Gly Leu
     47 atc gag cag gcg gcc aag cag gac gtg cgc ctg atc gca ttc cca gag
                                                                           144
     48 Ile Glu Gln Ala Ala Lys Gln Asp Val Arg Leu Ile Ala Phe Pro Glu
     51 act tgg att ccc ggc tat ccc ttt tgg ata tgg ctg ggc gcg ccg gct
                                                                           192
     52 Thr Trp Ile Pro Gly Tyr Pro Phe Trp Ile Trp Leu Gly Ala Pro Ala
             50
                                 55
     55 tgg ggc atg cgc ttc gtc cag cgc tat ttc gag aat tcg ctc gtg cgc
                                                                           240
     56 Trp Gly Met Arg Phe Val Gln Arg Tyr Phe Glu Asn Ser Leu Val Arg
     59 ggc agc aag cag tgg cag gcc ctg gcg gat gcg gcc cgc cgc cac ggc
                                                                           288
     60 Gly Ser Lys Gln Trp Gln Ala Leu Ala Asp Ala Ala Arg Arg His Gly
     61
                                              90
                         85
     63 atg cat gtc gtg gcc ggc tat agc gag cgc gcg ggc ggc agc ctc tat
                                                                           336
     64 Met His Val Val Ala Gly Tyr Ser Glu Arg Ala Gly Gly Ser Leu Tyr
                                         105
                    100
                                                                           384
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67 atg ggc cag gcg atc ttc ggc ccc gat ggc gat ctg atc gcc gcg cgc

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/751,299



Input Set : A:\DIVER1440-2final.txt
Output Set: C:\CRF3\05312001\1751299.raw

68 Met Gly Gln Ala Ile Phe Gly Pro Asp Gly Asp Leu Ile Ala Ala Arg 120 115 71 cgc aag ctc aag cct acc cat gcg gag cgc acc gtg ttc qqc qaq qqa 432 72 Arg Lys Leu Lys Pro Thr His Ala Glu Arg Thr Val Phe Gly Glu Gly 135 75 gac ggc agc cat ctc gcg gtg cac gat acc gcc atc ggg cgc ctc ggc 480 76 Asp Gly Ser His Leu Ala Val His Asp Thr Ala Ile Gly Arg Leu Gly 150 155 79 gcg ctc tgt tgc tgg gag cac atc cag cca ttg tcg aaa tac gcc atg 528 80 Ala Leu Cys Cys Trp Glu His Ile Gln Pro Leu Ser Lys Tyr Ala Met 165 83 tac gcc gcc gac gaa cag gtc cac gtc gcg tcg tgg ccg agc ttc agc 576 84 Tyr Ala Ala Asp Glu Gln Val His Val Ala Ser Trp Pro Ser Phe Ser 180 185 87 ctc tat cgc ggc atg gcc tat gcg ctc gga ccg gag gtc aat acc gcc 624 88 Leu Tyr Arg Gly Met Ala Tyr Ala Leu Gly Pro Glu Val Asn Thr Ala 195 200 91 gca age cag ate tae geg gte gag gge tge tae gtg etg geg teg 672 92 Ala Ser Gln Ile Tyr Ala Val Glu Gly Gly Cys Tyr Val Leu Ala Ser 215 95 tgc gcg acc gtt tcg ccg gag atg atc aag gta ttg gtg gat acg ccc 720 96 Cys Ala Thr Val Ser Pro Glu Met Ile Lys Val Leu Val Asp Thr Pro 230 235 99 gac aag gag atg ttc ctc aag gcc ggc ggt ggt ttt gcc atg att ttc 768 100 Asp Lys Glu Met Phe Leu Lys Ala Gly Gly Phe Ala Met Ile Phe 245 103 ggg ccc gac ggc cgc gcc ctg gcc gag ccg ctc ccg gag acc gaa gag 816 104 Gly Pro Asp Gly Arg Ala Leu Ala Glu Pro Leu Pro Glu Thr Glu Glu 105 260 107 gga ctg ctg gtc gcc gat atc gac ctc ggc atg atc gcg ttg gcc aag 864 108 Gly Leu Leu Val Ala Asp Ile Asp Leu Gly Met Ile Ala Leu Ala Lys 280 111 gcg gcg gcc gat ccg gcg ggc cac tat tca cgg ccc gac gta acg cgg 912 112 Ala Ala Ala Asp Pro Ala Gly His Tyr Ser Arg Pro Asp Val Thr Arg 295 115 ctg ctg ctg gat cga cgt ccg gcc caa cgc gtc gtc acg ctt gat gcc 960 116 Leu Leu Leu Asp Arg Pro Ala Gln Arg Val Val Thr Leu Asp Ala 310 119 gca ttc gaa ccg caa aac gag gac aag ggc gac gcg ccc gcg ctg cgc 1008 120 Ala Phe Glu Pro Gln Asn Glu Asp Lys Gly Asp Ala Pro Ala Leu Arg 325 123 gtg gtg gcg gaa agc gcc gcc gcc gcg cag tag´ 1041 124 Val Val Ala Glu Ser Ala Ala Ala Gln 340 128 <210> SEQ ID NO: 2 129 <211> LENGTH: 346 130 <212> TYPE: PRT 131 <del>≤213></del>--ORGANISM: Unknown Organism W--> 132 <220> FEATURE: -> See next page.



#### RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/751,299

DATE: 05/31/2001 TIME: 17:55:10

Input Set : A:\DIVER1440-2final.txt
Output Set: C:\CRF3\05312001\I751299.raw

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Input Set : A:\DIVER1440-2final.txt Output Set: C:\CRF3\05312001\I751299.raw

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	192	<220					•											
	193	<22	1> NA	AME/I	KEY:	CDS												
:	194	<222	2> LO	CAT:	ON:	(1)	(10	014)										
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			Lys	Glu	Ala		Lys	Val	Ala	Cys		Gln	Ala	Ala	Pro		Tyr	
	199	. 1				5					10					15		0.5
															atg			96
	202	мес	Asp	ьeu	20	Ата	THE	vai	Asp	цуs 25	Thr	тте	GIU	ьeu	Met 30	GIU	GIU	
		aca	aca	cat		22+	act	cat	cta		acc	+++	cca	maa	act	taa	a++	144
															Thr			723
	207			35				9	40				110	45		тър	110	
		cca	qqc		cca	tgg	ttt	ctt	tgg	ctt	gac	tca	cca	gca	tgg	qca	atq	192
															Trp			
	211		50			-		55	_		_		60		_			
															gat			240
			Phe	Val	Arg	Gln	_	His	Glu	Asn	Ser	Leu	Glu	Leu	Asp	Gly	Pro	
	215						70					75					80	
															atc			288
		GIn	Ala	Lys	Arg		Ser	Asp	Ата	Ala	_	Arg	Leu	GLY	Ile		Val	
	219	200	a+ ~	~~~	2+~	85	~~~	000	ata	aat	90	200	a++	+	a to	95	~~~	336
															atc Ile			336
	223	1111	пец	СТУ	100	Ser	GIU	Arg	vai	105	СТУ	1111	пец	TYL	110	261	GIII	
		taa	ttc	ata		gat	aat	aat	gac		att	aaa	acc	caa	cga	aaσ	tta	384
															Arg			
	227	-		115	-	-		-	120			_		125		•		
															gat			432
		Lys		Thr	Phe	Val	Glu	_	Thr	Leu	Phe	Gly	Glu	Gly	Asp	Gly	Ser	
	231		130					135					140					
															ggc			480
			Leu	Ala	Val	Phe		Thr	Ser	Val	Gly	-	Leu	Gly	Gly	Leu	_	
		145	<b>.</b>				150					155					160	E 2.0
															tat Tyr			528
	239	Суѕ	тър	GIU	птэ	165	GTII	PIO	Leu	1111	170	тÀт	АІА	ьец	ıyı	175	GIII	
		aat	maa	aaa	att		tat	aca	act	taa		agc	+++	agc	ctt		cct	576
															Leu			3,0
	243		<u> </u>	<u> </u>	180		<b>0 7 0</b>			185		~			190	- 1 -	<del>-</del> -	
		aat	gca	qca		gcc	cta	qaa	cct		gtc	aat	gta	gca	gcc	tct	cga	624
															Āla			
	247			195	-			-	200	-				205			-	
															tgt			672
2	250	Ile	Tyr	Ala	Val	Glu	Gly	Gln	Cys	Phe	Val	Leu	Ala	Ser	Cys	Ala	Leu	



Input Set : A:\DIVER1440-2final.txt
Output Set: C:\CRF3\05312001\I751299.raw

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210
                                 215
     251
     253 gtt tca caa tcc atg atc gat atg ctt tgt aca gat gac gaa aag cat
                                                                            720
     254 Val Ser Gln Ser Met Ile Asp Met Leu Cys Thr Asp Asp Glu Lys His
                             230
                                                  235
     257 gcg ttg ctt ctg gct ggt gga cac tca cgt atc ata ggg cct gat
                                                                            768
     258 Ala Leu Leu Leu Ala Gly Gly Gly His Ser Arg Ile Ile Gly Pro Asp
                         245
                                              250
     261 ggt ggt gac ttg gtc gcg cct ctt gcc gaa aat gaa gag ggt att ctc
     262 Gly Gly Asp Leu Val Ala Pro Leu Ala Glu Asn Glu Glu Gly Ile Leu
                     260
                                         265
     265 tac gca aac ctt gat cct gga gta cgc atc ctt gct aaa atg gcg gca
     266 Tyr Ala Asn Leu Asp Pro Gly Val Arg Ile Leu Ala Lys Met Ala Ala
                                     280
     269 gac cct gct ggt cat tat tcc cgt ccc gac att act cgc ttg cta ata
                                                                            912
     270 Asp Pro Ala Gly His Tyr Ser Arg Pro Asp Ile Thr Arg Leu Leu Ile
     271
             290
     273 gat cgc agc cct aaa tta ccg gta gtt gaa att gaa ggt gat ctt cgt
                                                                            960
     274 Asp Arg Ser Pro Lys Leu Pro Val Val Glu Ile Glu Gly Asp Leu Arg
                             310
                                                 315
     277 cct tac gct ttg ggt aaa gcg tct gag acg ggt gcg caa ctc gaa gaa
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     278 Pro Tyr Ala Leu Gly Lys Ala Ser Glu Thr Gly Ala Gln Leu Glu Glu
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                 35
                                      40
     299 Pro Gly Tyr Pro Trp Phe Leu Trp Leu Asp Ser Pro Ala Trp Ala Met
                                  55
     301 Gln Phe Val Arg Gln Tyr His Glu Asn Ser Leu Glu Leu Asp Gly Pro
                              70
     303 Gln Ala Lys Arg Ile Ser Asp Ala Ala Lys Arg Leu Gly Ile Met Val
     305 Thr Leu Gly Met Ser Glu Arg Val Gly Gly Thr Leu Tyr Ile Ser Gln
                     100
                                         105
     307 Trp Phe Ile Gly Asp Asn Gly Asp Thr Ile Gly Ala Arg Arg Lys Leu
                                    120
     309 Lys Pro Thr Phe Val Glu Arg Thr Leu Phe Gly Glu Gly Asp Gly Ser
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135

310

130

VERIFICATION SUMMARYDATE: 05/31/2001PATENT APPLICATION: US/09/751,299TIME: 17:55:11

Input Set : A:\DIVER1440-2final.txt
Output Set: C:\CRF3\05312001\I751299.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:132 M:258 W: Mandatory Feature missing, <220> FEATURE: L:289 M:258 W: Mandatory Feature missing, <220> FEATURE: